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Reviewer Comments:
<210>   3
<211>   17
<212>   PRT
<213>   Artificial Sequence


<220>
<223>   Synthetic construct


<220>
<221>   MISC_FEATURE
<222>   (1)..(1)
<223>   Xaa is Asp or absent


<220>
<221>   MISC_FEATURE
<222>   (2)..(2)
<223>   Xaa is Arg or absent


<220>
<221>   MISC_FEATURE
<222>   (3)..(3)
<223>   Xaa is Met or absent


<220>
<221>   MISC_FEATURE
<222>   (4)..(4)
<223>   Xaa is Pro or absent


<220>
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<221>   MISC_FEATURE
<222>   (6)..(6)
<223>   Xaa is Arg or Lys

<220>
<221>   MISC_FEATURE
<222>   (14)..(14)
<223>   Xaa is Ser or Thr

<220>
<221>   MISC_FEATURE
<222>   (17)..(17)
<223>   Xaa is Lsy or absent

<400>   3

Xaa Xaa Xaa Xaa Cys Xaa Asn Phe Phe Trp Lys Thr Phe Xaa Ser Cys
1   5   10   15

Xaa

The above <223> explanation for the "Xaa" at location 17 contains an
error: "Lsy" is not a valid amino acid designator.
****

Application No: 10533115   Version No: 3.0

Input Set:

Output Set:

Started: 2008-09-20 07:30:53.095
Finished: 2008-09-20 07:30:55.585
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 490 ms
Total Warnings: 3
Total Errors: 0
No. of SeqIDs Defined: 3
Actual SeqID Count: 3

Error code   Error Description

W   213   Artificial or Unknown found in <213> in SEQ ID (1)

W   213   Artificial or Unknown found in <213> in SEQ ID (2)

W   213   Artificial or Unknown found in <213> in SEQ ID (3)

<210> 1
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic construct

<400> 1
tggccgtctt cattttctgc tcgccgctca ctttgaccaa g   41

<210> 2
<211> 42
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic construct

<400> 2
gtgggcaaga tgcgcgctgt gaatggggtt ggcgcagctg tt   42

<210> 3
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic construct

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa is Asp or absent

<220>
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<222> (17)..(17)
<223> Xaa is Lsy or absent

<400> 3

Xaa Xaa Xaa Xaa Cys Xaa Asn Phe Phe Trp Lys Thr Phe Xaa Ser Cys
1   5   10   15

Xaa